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## SEQUENCE LISTING

Oldham, Elizabeth R. Soto, Hortensia Liu, Ying Hudak, Susan A. Homey, Bernhard Morales, Janine M. Kellerman, Sirid-Aimee McEvoy, Leslie M. Bowman, Edward P. Zlotnik, Albert

## <120> CHEMOKINE AND RECEPTOR USES; COMPOSITIONS; METHODS <130> DX0882XK <140> US09/898,751 <141> 2001-07-02 <150> US09/471,549 <151> 1999-12-23 <150> US60/136,570 <151> 1999-05-27 <150> US60/113,858 <151> 1998-12-24 <160> 16 <170> PatentIn version 3.1 <210> 1 <211> 1089 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(1086) <223> <400> 1 atg ggg acg gag gtt tta gag cag gtt tcc tgg ggc cat tac tct ggg 48 Met Gly Thr Glu Val Leu Glu Gln Val Ser Trp Gly His Tyr Ser Gly gat gaa gag gac gca tac tog got gag coa ctg cog gag ott tgo tac 96 Asp Glu Glu Asp Ala Tyr Ser Ala Glu Pro Leu Pro Glu Leu Cys Tyr 20 aag god gat gto dag god tto ago ogg god tto daa dod agt gto tod 144 Lys Ala Asp Val Gln Ala Phe Ser Arg Ala Phe Gln Pro Ser Val Ser 35 ctg acg ctg gct gcg ctg ggt ctg gcc ggc aat ggc ctg gtc ctg gcc

Leu Thr Leu Ala Ala Leu Gly Leu Ala Gly Asn Gly Leu Val Leu Ala

50

192

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cgc cgc ggc Arg Arg Gly	tgc ccc c Cys Pro A 340	ge egg eed rg Arg Pro	c cgc ctt Arg Leu 345	tct tcc Ser Ser	tgc tca Cys Ser 350	gct ccc Ala Pro	1056
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Lys Ala Asp 35	Val Gln A	la Phe Sei 40	r Arg Ala	Phe Gln	Pro Ser 45	Val Ser	
Leu Thr Leu 50	Ala Ala L	eu Gly Let 55	ı Ala Gly	Asn Gly 60	Leu Val	Leu Ala	
Thr His Leu 65	Ala Ala A		a Ala Arg	Ser Pro 75	Thr Ser	Ala His 80	
Leu Leu Gln	Leu Ala L 85	eu Ala Asp	p Leu Leu 90	Leu Ala	Leu Thr	Leu Pro 95	
Phe Ala Ala	Ala Gly A 100	la Leu Gli	o Gly Trp 105	Ser Leu	Gly Ser 110	Ala Thr	
Cys Arg Thr 115	Ile Ser G	ly Leu Tyr 120		Ser Phe	His Ala 125	Gly Phe	
Leu Phe Leu 130	Ala Cys I	le Ser Ala 135	a Asp Arg	Tyr Val 140	Ala Ile	Ala Arg	
Ala Leu Pro 145		ro Arg Pro 50	Ser Thr	Pro Gly 155	Arg Ala	His Leu 160	

Val Ser Val Ile Val Trp Leu Leu Ser Leu Leu Leu Ala Leu Pro Ala Leu Leu Phe Ser Gln Asp Gly Gln Arg Glu Gly Gln Arg Arg Cys Arg Leu Ile Phe Pro Glu Gly Leu Thr Gln Thr Val Lys Gly Ala Ser Ala 200 Val Ala Gln Val Ala Leu Gly Phe Ala Leu Pro Leu Gly Val Met Val 215 Ala Cys Tyr Ala Leu Leu Gly Arg Thr Leu Leu Ala Ala Arg Gly Pro 240 Glu Arg Arg Arg Ala Leu Arg Val Val Ala Leu Val Ala Ala Phe Val Val Leu Gln Leu Pro Tyr Ser Leu Ala Leu Leu Asp Thr Ala 260 Asp Leu Leu Ala Ala Arg Glu Arg Ser Cys Pro Ala Ser Lys Arg Lys 280 Asp Val Ala Leu Leu Val Thr Ser Gly Leu Ala Leu Ala Arg Cys Gly 295 Leu Asn Pro Val Leu Tyr Ala Phe Leu Gly Leu Arg Phe Arg Gln Asp 315 310 Leu Arg Arg Leu Leu Arg Gly Gly Ser Ser Pro Ser Gly Pro Gln Pro 330 325 Arg Arg Gly Cys Pro Arg Arg Pro Arg Leu Ser Ser Cys Ser Ala Pro 345 Thr Glu Thr His Ser Leu Ser Trp Asp Asn 360 355 <210> 3

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aag Lys	gct Ala	gat Asp 35	gtc Val	cag Gln	gct Ala	ttc Phe	agt Ser 40	cgg Arg	gcc Ala	ttc Phe	caa Gln	ccc Pro 45	agt Ser	gtc Val	tcc Ser	144
ctg Leu	atg Met 50	gtg Val	gct Ala	gta Val	ctg Leu	ggt Gly 55	ctg Leu	gct Ala	ggc Gly	aat Asn	ggc Gly 60	cta Leu	gtc Val	ttg Leu	gcc Ala	192
acc Thr 65	cat His	ctg Leu	gca Ala	gcc Ala	aga Arg 70	cga Arg	act Thr	acc Thr	cga Arg	tct Ser 75	ccc Pro	acc Thr	tcc Ser	gtt Val	cac His 80	240
ctg Leu	ctc Leu	cag Gln	ttg Leu	gcc Ala 85	ctg Leu	gct Ala	gac Asp	ctt Leu	tta Leu 90	ttg Leu	gcc Ala	ctg Leu	act Thr	ttg Leu 95	cct Pro	288
ttt Phe	gct Ala	gca Ala	gca Ala 100	ggg Gly	gct Ala	ctt Leu	cag Gln	ggc Gly 105	tgg Trp	aat Asn	cta Leu	gga Gly	agt Ser 110	acc Thr	acc Thr	336
tgc Cys	cgt Arg	gcc Ala 115	atc Ile	tca Ser	ggc Gly	ctc Leu	tac Tyr 120	tcg Ser	gcc Ala	tct Ser	ttc Phe	cac His 125	gct Ala	ggc Gly	ttc Phe	384
ctc Leu	ttc Phe 130	cta Leu	gcc Ala	tgt Cys	atc Ile	agc Ser 135	gcc Ala	gac Asp	cgc Arg	tat Tyr	gtg Val 140	gcc Ala	atc Ile	gca Ala	cga Arg	432
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gtg Val	gcg Ala 210	Gln	gtg Val	gtc Val	ctc Leu	ggc Gly 215	Phe	gcg Ala	ctc Leu	cct Pro	ctg Leu 220	GIY	gtc Val	atg Met	gca Ala	672

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Ala Cys Tyr Ala Leu Leu Gly Arg Thr Leu Leu Ala Ala Arg Gly Pro 225 230 235 240	
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gat cta ctg gca gcc cgc gag cgg agc tgc tcc tcc agc aag cgc aag Asp Leu Ala Ala Arg Glu Arg Ser Cys Ser Ser Ser Lys Arg Lys 275 280 285	864
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Leu Met Val Ala Val Leu Gly Leu Ala Gly Asn Gly Leu Val Leu Ala 50 55 60	

Thr His Leu Ala Ala Arg Arg Thr Thr Arg Ser Pro Thr Ser Val His

Leu Leu Gln Leu Ala Leu Ala Asp Leu Leu Leu Ala Leu Thr Leu Pro 85 90 95

Phe Ala Ala Gly Ala Leu Gln Gly Trp Asn Leu Gly Ser Thr Thr
100 105 110

Cys Arg Ala Ile Ser Gly Leu Tyr Ser Ala Ser Phe His Ala Gly Phe 115 120 125

Leu Phe Leu Ala Cys Ile Ser Ala Asp Arg Tyr Val Ala Ile Ala Arg 130 135 140

Ala Leu Pro Ala Gly Gln Arg Pro Ser Thr Pro Ser Arg Ala His Leu 145 150 155 160

Val Ser Val Phe Val Trp Leu Leu Ala Leu Phe Leu Ala Leu Pro Ala 165 170 175

Leu Leu Phe Ser Arg Asp Gly Pro Arg Glu Gly Gln Arg Arg Cys Arg 180 185 190

Leu Ile Phe Pro Glu Ser Leu Thr Gln Thr Val Lys Gly Ala Ser Ala 195 200 205

Val Ala Gln Val Val Leu Gly Phe Ala Leu Pro Leu Gly Val Met Ala 210 215 220

Ala Cys Tyr Ala Leu Leu Gly Arg Thr Leu Leu Ala Ala Arg Gly Pro 225 230 235 240

Glu Arg Arg Arg Ala Leu Arg Val Val Val Ala Leu Val Val Ala Phe
245 250 255

Val Val Leu Gln Leu Pro Tyr Ser Leu Ala Leu Leu Leu Asp Thr Ala 260 265 270

Asp Leu Leu Ala Ala Arg Glu Arg Ser Cys Ser Ser Ser Lys Arg Lys 275 280 285

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45

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														agg Arg 90		394
			aaa Lys 95													436
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ttgt	cct	gca t	acto	gatga	aa a	ctact	gat	g tev	/gctg	ggtc	tgaa	aagga	acc 1	tacca	agaagc	556
taaa	atct	cca a	agaat	gcca	at t	cect	atc	cta	aatga	attc	aato	ctcc	ctt a	accct	gacca	616
atca	agtg	gee (	caaat	tttc	cc ag	geee	cttgo	cto	ccag	gaac	ccca	agcco	cag a	aacto	cttcag	676
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Leu	His -5	Ala	Ser	Glu	Ala -1	Ile 1	Leu	Pro	Ile	Ala 5	Ser	Ser	Cys	Сув	Thr 10	
Glu	Val	Ser	His	His 15	Ile	Ser	Arg	Arg	Leu 20	Leu	Glu	Arg	Val	Asn 25	Met	
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Leu	His	Val 45	Lys	Arg	Arg	Arg	Ile 50	Cys	Val	Ser	Pro	His 55	Asn	His	Thr	
Val	Lys 60	Gln	Trp	Met	Lys	Val 65	Gln	Ala	Ala	Lys	Lys 70	Asn	Gly	Lys	Gly	

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agg gag cgg agt Arg Glu Arg Ser 35	ccg atg tct Pro Met Ser	cca aca agc Pro Thr Ser 40	cag aga cta agt Gln Arg Leu Ser 45	ctg gaa 144 Leu Glu
gcc ccc agc ctc Ala Pro Ser Leu 50	cca ctg aga Pro Leu Arg 55	agc tgg cat Ser Trp His	ccg tgg aac aag Pro Trp Asn Lys 60	act aag 192 Thr Lys
cag aag caa gaa Gln Lys Gln Glu 65	gcc ttg cct Ala Leu Pro 70	Leu Pro Ser	agc act agc tgc Ser Thr Ser Cys 75	tgt act 240 Cys Thr 80
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cac atg gaa ctg His Met Glu Leu 100	cag gag gcc Gln Glu Ala	gat ggg gac Asp Gly Asp 105	tgt cac ctc cag Cys His Leu Gln 110	gct gtc 336 Ala Val
gtg ctt cac ctg Val Leu His Leu 115	Ala Arg Arg	Ser Val Cys 120	Val His Pro Gin 125	Asn Arg
Ser Leu Ala Arg 130	Trp Leu Glu 135	Arg Gln Gly	aaa agg ctc caa Lys Arg Leu Gln 140	Gly Thr
Val Pro Ser Leu 145	Asn Leu Val 150	Leu Gln Lys	aaa atg tac tca Lys Met Tyr Ser 155	160
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Ala Pro Ser Leu Pro Leu Arg Ser Trp His Pro Trp Asn Lys Thr Lys 55

Gln Lys Gln Glu Ala Leu Pro Leu Pro Ser Ser Thr Ser Cys Cys Thr 80 70

Gln Leu Tyr Arg Gln Pro Leu Pro Ser Arg Leu Leu Arg Arg Ile Val 90 85

His Met Glu Leu Gln Glu Ala Asp Gly Asp Cys His Leu Gln Ala Val 105

Val Leu His Leu Ala Arg Arg Ser Val Cys Val His Pro Gln Asn Arg 120

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Glu	Val	Ser	His	His 15	Val	Ser	Gly	Arg	Leu 20	Leu	Glu	Arg	Val	Ser 25	Ser	
Cys	Ser	Ile	Gln 30	Arg	Ala	Asp	Gly	Asp 35	Cys	Asp	Leu	Ala	Ala 40	Val	Ile	
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Asn 75	Val	Сув	Ser	Gly	Lys 80	Lys	Gln	Pro	Ser	Arg 85	Lys	qsA	Arg	Lys	Gly 90	
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Gln Asn Pr	Ser Leu 60	Ser Gln	Trp Phe 65	Glu His	Gln Glu	Arg Lys	: Leu
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Ser Cys Cys Thr Gln Leu Tyr Arg Gln Pro Leu Pro Ser Arg Leu Leu Arg Arg Ile Val His Met Glu Leu Gln Glu Ala Asp Gly Asp Cys His 30 Leu Gln Ala Val Val Leu His Leu Ala Arg Arg Ser Val Cys Val His Pro Gln Asn Arg Ser Leu Ala Arg Trp Leu Glu Arg Gln Gly Lys Arg Leu Gln Gly Thr Val Pro Ser Leu Asn Leu Val Leu Gln Lys Lys Met Tyr Ser Asn Pro Gln Gln Asn <210> 15 <211> 32 <212> DNA <213> Homo sapiens <400> 15 32 atotggcacc acaccttcta caatgagctg cg <210> 16 <211> 32 <212> DNA <213> Homo sapiens <400> 16 cgtcatactc ctgcttgctg atccacatct gc 32